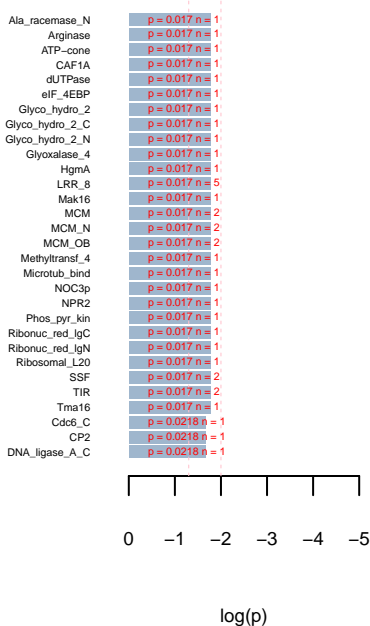


Oecy.Fem-Mal_difpos domain



n = 72 genes | p adjust BH TRUE